REMARKS

Attached hereto is a marked-up version of the changes made to the specification and

claims by the current amendment. The attached page is captioned "VERSION WITH

MARKINGS TO SHOW CHANGES MADE".

Applicant has amended the specification and abstract to comport with United States

Patent and Trademark Office Rules. No new matter is added. Applicant has amended the claims

to eliminate multiple dependency and to comport with U.S. practice, which is totally unrelated to

patentability. No new matter is added.

In view of the above, it is respectfully believed that all the presently submitted claims are

allowable and a Formal Notice of Allowance is courteously solicited. It is believed that the

application is in condition for allowance, however, if the Examiner feels otherwise, a telephone

interview is respectfully requested. An early notice of allowance is solicited.

Respectfully submitted,

Date: 3/18/07

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VERSION WITH MARKINGS TO SHOW CHANGES MADE

Following is a marked-up version of the Specification with all changes shown by conventional comparison (underlining and bracketing):

[TITLE OF THE INVENTION]

LABELED NEUROTENSIN DERIVATIVES

APPLICATION CROSS-REFERENCES

This application claims priority of International Application No. PCT/US00/17509, filed June 22, 2000. This application also claims priority of U.S. Provisional Application No. 60/140,913, filed June 24, 1999 and U.S. Provisional Application No. 60/213,068, filed June 21, 2000.

[FIELD OF THE INVENTION] FIELD OF THE INVENTION

The present invention relates to: labeled peptide compounds, a method of preparing the compounds such that they are resistant to enzymatic degradation, a pharmaceutical composition comprising these compounds, and use of these compounds for the purposes of diagnosis and therapy.

[BACKGROUND OF THE INVENTION] BACKGROUND OF THE INVENTION

Neurotensin (NT) is a thirteen amino acid peptide, isolated from bovine hypothalamus and has the following structure: pGlu-Leu-Tyr-Glu-Asn-Lys-Pro-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:1) wherein pGlu is pyroglutamate. High concentrations of neurotensin receptors are found in discrete regions of the mammalian central nervous system including the brain and in the gut. In addition, neurotensin receptors are found in several tumor cells, including small cell lung carcinoma, exocrine pancreatic cancer (Reubi et al., 1998), Ewing sarcoma, meningiomas, medulloblastomas and astrocytomas. Normal pancreatic tissue and tissue from patients with pancreatitis or endocrine pancreatic cancer do not express neurotensin receptors (Reubi et al., 1998). It is estimated that there are 58,000 cases of exocrine pancreatic cancer per year in the United States and Europe. The five year survival rate for patients with exocrine pancreatic cancer is low, in the range of 5-10%. Current diagnosis for this cancer uses a combination of radiologic procedures and biopsies. An early diagnostic method coupled with a therapeutic counterpart may have a profound effect on survival and quality of life.

Structure-activity relationships have shown that the C-terminal sequence (amino acid residues 8-13 (named NT(8-13))) of the natural neurotensin is sufficient for preserving high affinity receptor binding (Granier et al., 1982; Kitabgi et al., 1985). The affinity of this analog is comparable to that of natural neurotensin in two different binding assays, i.e., the binding assay on rat brain synaptic membranes and on HT 29 cells which express neurotensin receptors.

Unfortunately this truncated peptide has poor *in vitro* stability. One site of enzymatic instability is the Arg⁸-Arg⁹ bond which has a serum t₂ of 5 minutes. The serum stability was increased with the Lys⁸(Ψ-CH₂NH)Arg⁹ pseudo-peptide DTPA-Lys⁸(Ψ-CH₂NH)Arg⁹-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:2) (Tourwe et al., 1998).

Two radioiodinated derivatives of neurotensin are described in the literature. Because there are two tyrosine residues, iodination yields a complex mixture of products. These derivatives are difficult to purify and each one possesses different biological properties. To overcome this problem, Mazella et al. (1983) synthesized monoiodo-125-[Trp 11]-neurotensin. This derivative has a K_d of 0.1 nM binding to rat brain synaptic membranes. The same group of researchers later succeeded in preparing the monoiodo-125 derivative of natural neurotensin derivative. This radioiodo analog has a K_d of 0.26 nM for binding on human brain neurotensin membranes.

These iodinated derivatives of natural neurotensin peptides are unsuitable for imaging and therapy of tumors expressing neurotensin receptors because of difficulty in the method of preparation as well as the instability of these derivatives. The instability results from rapid deiodination and also from the enzymatic degradation of the natural neurotensin peptide bonds.

NT(8-13) contains only one Tyr residue which can be selectively radioiodinated. Structure activity studies indicated that the iodination resulted in the loss of binding affinity by a factor of 20.

Since the early work, other radiolabeled neurotensin analogs have been prepared. Tourwe et al. (1998) have prepared diethylenetriamine pentaacetic acid (DTPA)-NT(8-13) (DTPA-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:3)) and found that the derivative had an affinity of 6.5 nM to HT 29 colon adenocarcinoma cells. The low tumor uptake in nude mice HT29 tumor was ascribed to the rapid metabolism of the DTPA-NT(8-13). The *in vivo* half-life

of neurotensin is less than 1.5 minutes and the major cleavage site has been shown to be the Arg^8 - Arg^9 bond (Lee et al., 1984; Aronin et al., 1982). Hence, neurotensin analogs in which the peptide bonds were sequentially replaced by $\Psi(CH_2NH)$ were prepared and a large drop in affinity was observed. The compounds DTPA-Lys⁸- $\Psi(CH_2NH)$ - Arg^8 -NT(8-13) and DTPA-Lys⁸- $\Psi(CH_2NH)$ -Lys⁸-NT(8-13) had a K_d of 13 and 7.4 nM, respectively.

An analysis of the above compounds indicated that the stability of the compounds in the serum is not sufficient for these compounds to be used as radiolabeled imaging and therapeutic agents.

The publications and other materials used herein to illuminate the background of the invention or provide additional details respecting the practice, are incorporated by reference, and for convenience are respectively grouped in the appended List of References.

[SUMMARY OF THE INVENTION] SUMMARY OF THE INVENTION

Neurotensin analogs containing Arg mimics have been synthesized. Studies demonstrate that replacement of Arg⁸ does not significantly affect the binding affinity and replacement of Arg⁹ is not tolerated. The best results are obtained by replacing Arg⁸ with (4-Gu)Phe or Gly(PipAm) ((N-amidinopiperidinyl)glycine) as the arginine surrogate. The IC₅₀ values for these two peptides are comparable to native neurotensin. While serum stability is improved by the incorporation of Arg mimics, optimum stability is not achieved by that change alone. Another source of instability is the C-terminus with the Ile-Leu-OH being metabolized. Replacement of the C-terminus with a bulkier side chain stabilized the bond from degradation. Replacement of Ile with tBuGly results in no loss of binding affinity, although the presence of the pseudo-peptide bond or the C-terminal amide abolishes receptor affinity. Neurotensin analogs are described which are useful for diagnostic and therapeutic purposes.

[DETAILED DESCRIPTION OF THE INVENTION] DETAILED DESCRIPTION OF THE INVENTION

Neurotensin has a short half-life *in vivo* thereby limiting its usefulness as a diagnostic or therapeutic agent. It is desirable to have a compound with a stability such that 70-80% of the

injected compound is present in the subject=s serum and urine at the end of 4 hours after administration. Several neurotensin analogs have been synthesized, as disclosed herein, which are stable and still show strong binding affinity to neurotensin receptors.

The phrase "spacer unit" as used herein designates any combination of amino acids or amino acid residues, a combination of amino acids or amino acid residues with a non-amino acid moiety, or any non-amino acid moiety which removes (i.e., spaces) a chelating moiety from a binding portion of a peptide.

The phrase "selective affinity" as used herein means a binding affinity at least in the micromolar or stronger binding. Selective affinity includes a K_d in the micromolar, nanomolar or stronger range.

The initial changes made to neurotensin consisted of replacing the arginine at amino acid position 1 of NT(8-13) with an arginine mimic. It was further recognized that there is a secondary site of metabolism between Ile-Leu-OH (AA₅-AA₆). Hence Ile was replaced by a bulkier side chain to stabilize the bond from further degradation. The combination of both modifications is essential for the stabilization of natural neurotensin derivatives as well as its analogs. Other modifications such as replacement of Arg-9-neurotensin (AA₂) results in the loss of binding affinity.

The present invention is further detailed in the following Examples, which are offered by way of illustration and are not intended to limit the invention in any manner. Standard techniques well known in the art or the techniques specifically described below are utilized.

Example 1

Peptide Synthesis

Solid phase peptide synthesis (SPPS) was performed using an Applied Biosystems Model 432A "Synergy" Peptide synthesizer employing Fmoc (9-fluorenylmethoxycarbonyl) strategy. Instrument protocol required 25 µmol of starting resin and 75 µmol of subsequent Fmocprotected amino acids activated by a combination of *N*-hydroxy-benzotriazole (HOBt) and 2-(1-H Benzotriazol-1-yl)-1,1,3,3-tetramethyluroniumhexafluorophosphate (HBTU). Tri-t-butyl DTPA (75 µmol), prepared internally, was placed in an "amino acid column" at the appropriate

location. The Fmoc-protected amino acids were purchased commercially unless otherwise stated; the pre-packaged amino acids were obtained from PE Biosystems while those unavailable in pre-packed cartridges, such as the *D* amino acids, were supplied by BACHEM or Novabiochem. The arginine mimics were purchased from RSP Amino Acid Analogues. All other peptide synthesis reagents were obtained from PE Biosystems. Cleavage and deprotection were accomplished using 85% TFA/5% thioanisole/5% phenol/5% water. The TFA was supplied by Pierce Chemical while the other cleavage reagents were purchased from Aldrich. The crude peptide was isolated by precipitation with t-butyl methyl ether (Sigma) and purified by reverse phase HPLC using an acetonitrile/water gradient containing 0.1% TFA. Molecular weight determination was accomplished by mass spectrometry operating in the electrospray mode (ESI).

Example 2

Competitive Binding Assay

Frozen tissue sections from receptor-positive human tumors were used in a competitive binding assay. Tissue samples were incubated with radioiodinated native neurotensin for 150 minutes at room temperature. Increasing amounts of the cold neurotensin derivatives were then added to generate competitive inhibition curves from which the IC₅₀ values were extrapolated.

Example 3

Standard Labeling Protocol

All reagents were purchased from Sigma unless otherwise noted. The peptide and ¹¹¹InCl₃ in 0.05 N HCl were incubated in a sodium acetate/ascorbic acid buffer for 30 minutes at room temperature. The reaction was diluted with 5% ethanol/95% PBS before using in subsequent *in vitro* assays.

Example 4

In Vitro Serum and Urine Stability Assays

Human serum was purchased from Sigma (H 1388). The labeled peptide was incubated in either human serum or rat urine at 37EC for the specified amount of time. An aliquot was filtered through a 0.45 μm syringe filter prior to HPLC injection.

Example 5

Arginine Mimics

An arginine mimic consists of a glycine moiety connected to a guanidino group with a spacer. The preferred ones have the formula:

 $\label{eq:cm-R3-(CA)_n-AA_1-AA_2-AA_3-AA_4-AA_5-AA_6-OH,} CM-R_3-(CA)_n-AA_1-AA_2-AA_3-AA_4-AA_5-AA_6-OH,$ wherein

CM is a chelating moiety or metal binding site wherein the chelating moiety is labeled with a metal isotope selected from ^{99m}Tc, ²⁰³Pb, ⁶⁷Ga, ¹¹¹In, ⁹⁷Ru, ⁶²Cu, ⁶⁴Cu, ¹⁸⁶Re, ¹⁸⁸Re, ⁹⁰Y, ¹²¹Sn, ¹⁶¹Tb, ¹⁵³Sm, ¹⁶⁶Ho, ¹⁰⁵Rh, ¹⁷⁷Lu or a radioactive halogen isotope on the understanding that

- i) if the label is a metal isotope, CM represents a chelating group suitable for the metal and
- ii) if the label is a radioactive halogen isotope, the halogen is attached to an aromatic ring, wherein the CM is attached directly or through a spacing group to the peptide, said chelating moiety being attached to the amine through an amide or urea bond or by any other modification which allows attachment of a chelate and which modifications are known to those of skill in the art,

wherein the chelating group is preferably derived from ethylene diamine tetraacetic acid (EDTA), diethylene triamine pentaacetic acid (DTPA), cyclohexyl 1,2-diamine tetraacetic acid (CDTA), ethyleneglycol-O,O=-bis(2-aminoethyl)-N,N,N',N'-diacetic acid (HBED), triethylene tetraamine hexaacetic acid (TTHA), 1,4,7,10-tetraazacyclododecane-N,N',N",N"'-tetraacetic acid (DOTA), 1,4,7-triazacyclononane-N,N',N"-triacetic acid (NOTA), 1,4,8,11-tetraazacyclotetradecane-N,N',N",N"'-tetraacetic acid (TETA) or a compound with a general formula

$$Y'''$$
 R_1 -N N-R₂
 Y'''
 S
 X
 PG
 Z

wherein

PG is a sulfur protecting group selected from alkanoyl, arylcarbonyl, arylalkanoyl, acetamidomethyl, tetrahydropyranyl and tetrahydrofuranyl,

Y', Y", and Y" are hydrogen or oxygen with the proviso that at least one of them is an O, R_1 and R_2 are hydrogen or alkyl (C_1 - C_3),

X = NH or S with the proviso that Y''' is hydrogen when X is S,

Z is PG if X is S, and

Z is hydroxyalkyl, aminoalkyl or carboxyalkyl;

 R_3 is DLys, DPhe or other D-amino acid, a spacer unit such as Gly-Gly-Gly, Gly-Ser-Gly, Tyr-Glu-Asn, DTyr-Glu-Asn, Phe-Glu-Asn or DPhe-Glu-Asn, or piperidinyl glycine (PipGly), aminomethylcyclohexylalanine (Amcha), other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring;

CA is a cyclic amino acid selected from Pro, Hyp, 4-oxo-proline [4OPro], pipecolic acid (PipCA), azetidinecarboxylic acid (AzeCA), other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring;

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$$n = 0, 1 \text{ or } 2;$$

 AA_1 = an amino acid containing a guanidino group except arginine, wherein the configuration at the α -carbon is either L- or D-, e.g.,

-HN
$$\alpha$$
-CO-
-HN α -N-
-HN

$$m = 0-6$$
;

R₄ is a cycloalkyl group (C₃-C₁₀), phenyl group, aralkyl group, substituted phenyl group or substituted aralkyl group with electron withdrawing or electron donating group with the proviso that the guanidino group is present at a position not occupied by the substituent on the phenyl group;

$$p = 1-7;$$

$$q = 1-7;$$

AA₂ is arginine, lysine, piperidinylglycine (PipGly), or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-configuration at the α -carbon, or AA₂ is an amino acid containing a guanidino group wherein the configuration at the α -carbon is either L- or D-, e.g.,

-HN
$$\alpha$$
-CO-
-HN α -N-
-HN

AA₃ is a cyclic amino acid selected from Pro, Hyp, 4-oxo-proline [4OPro], pipecolic acid (PipCA), azetidinecarboxylic acid (AzeCA), or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-configuration at the α -carbon;

AA₄ is Phe, Tyr, an isomer of Tyr, polyhydroxylated Phe, or other aromatic amino acid, wherein the amino acid can have the L- or D-configuration at the α -carbon;

AA₅ is Ile; and

AA₆ is Leu.

Preferred compounds include compounds I-VIII:

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

$$N$$
 H_2N
III

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

$$V$$
 N
 NH_2

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

VII

ΝΉ

Substitution of arginine with an arginine mimic or a constrained arginine as shown in compounds I-VIII increases the serum stability of these compounds considerably as shown in Table 1.

Table 1

Commound	v	% of compound present at 4 hours		
Compound	K _d	in serum	in urine	
DTPA-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:3)	40	1.6	13.0	
Compound I	8.6	14.1	1.5	
Compound II	2.7	19.5	16.6	

Compound III	68	4.3	6.0

As can be seen in Table 1, the presence of constrained arginine increased the serum stability considerably compared to the compound containing the Arg-Arg bond.

Novel neurotensin derivatives were prepared by replacing one or both arginines with the following mimics.

The binding affinities of the neurotensin derivatives with the incorporated Arg mimics are as follows:

	<u>K_d (nM)</u>
pGlu-Leu-Tyr-Glu-Asn-Lys-Pro-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:1)	3
DTPA-DLys-Pro-Arg-Phe(4-guanyl)-Pro-Tyr-Ile-Leu-OH	83
DTPA-DLys-Pro-Phe(4-guanyl)-Arg-Pro-Tyr-Ile-Leu-OH	8.6
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DTPA-DLys-Pro-Phe(4-guanyl)-Phe(4-guanyl)-Pro-Tyr-Ile-Leu-OH	175
DTPA-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:3)	40
DTPA-DLys-Pro-Arg-Aba(Apy)-Pro-Tyr-Ile-Leu-OH	1200
DTPA-DLys-Pro-Aba(Apy)-Arg-Pro-Tyr-Ile-Leu-OH	68
DTPA-DLys-Pro-Aba(Apy)-Aba(Apy)-Pro-Tyr-Ile-Leu-OH	>10000
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Ile-Leu-OH	2.7

<u>Table 2</u> <u>HPLC Evaluation of Serum and Urine Stability at 37EC</u>

Compound	Prep	Serum, t=0	Serum, t=4h	Urine,	Urine, t=4h
In-111-DTPA-R-R-P-Y-I-L	97.6%	92%	1.6%	96.7%	13.0%
(SEQ ID NO:3)	rt=14.2m	rt=14.2m	rt=14.2m	rt=14.2m	rt=14.2m
In-111-DTPA-DK-P-R-	95.5%	94%	16.3%	90.4%	0%
F(Gu)-P-Y-I-L	rt=15.1m	rt=15.1m	rt=15.1m	rt=14.7m	rt=14.7m
In-111-DTPA-DK-P-F(Gu)-	98.7%	98.7%	14.1%	94.1%	1.5%
R-P-Y-I-L	rt=14.6m	rt=14.6m	rt=14.6m	rt=14.6m	rt=14.6m
In-111-DTPA-DK-P-F(Gu)-	99.4%	99.0%	25.2%	92.3%	2.3%
F(Gu)-P-Y-I-L	rt=14.6m	rt=14.6m	rt=14.6m	rt=14.6m	rt=14.6m
In-111-DTPA-DK-P-R-	99.6%	98.1%	15.7%	96.7%	4.7%
Aba(Apy)-P-Y-I-L	rt=14.8m	rt=14.8m	rt=14.8m	rt=14.8m	rt=14.8m
In-111-DTPA-DK-P-	99.8%	96.9%	4.3%	95.6%	6.0%
Aba(Apy)-R-P-Y-I-L	rt=14.8m	rt=14.8m	rt=14.8m	rt=14.8m	rt=14.8m

In-111-DTPA-DK-P-	99.3%	Not	19.5%	Not	16.6%
G(PipAm)-R-P-Y-I-L	rt=13.8m	available	rt=13.8m	available	rt=13.8m

Example 6

Neurotensin Derivatives Containing Both Arg Mimics and C-Terminus Modifications

It was further recognized that there is a secondary site of metabolism between Ile-Leu-OH (AA₅-AA₆). Hence Ile was replaced by a bulkier side chain to stabilize the bond from further degradation. Compounds were made of the formula

$$\label{eq:cm-R3-(CA)_n-AA_1-AA_2-AA_3-AA_4-AA_5-AA_6-OH}$$
 wherein

CM is a chelating moiety or metal binding site wherein the chelating moiety is labeled with a metal isotope selected from ^{99m}Tc, ²⁰³Pb, ⁶⁷Ga, ¹¹¹In, ⁹⁷Ru, ⁶²Cu, ⁶⁴Cu, ¹⁸⁶Re, ¹⁸⁸Re, ⁹⁰Y, ¹²¹Sn, ¹⁶¹Tb, ¹⁵³Sm, ¹⁶⁶Ho, ¹⁰⁵Rh, ¹⁷⁷Lu or a radioactive halogen isotope on the understanding that

- i) if the label is a metal isotope, CM represents a chelating group suitable for the metal and
- ii) if the label is a radioactive halogen isotope, the halogen is attached to an aromatic ring, wherein the CM is attached directly or through a spacing group to the peptide, said chelating moiety being attached to the amine through an amide or urea bond or by any other modification which allows attachment of a chelate and which modifications are known to those of skill in the art,

wherein the chelating group is preferably derived from ethylene diamine tetraacetic acid (EDTA), diethylene triamine pentaacetic acid (DTPA), cyclohexyl 1,2-diamine tetraacetic acid (CDTA), ethyleneglycol-O,O=-bis(2-aminoethyl)-N,N,N',N'-diacetic acid (HBED), triethylene tetraamine hexaacetic acid (TTHA), 1,4,7,10-tetraazacyclododecane-N,N',N",N"'-tetraacetic acid (DOTA), 1,4,7-triazacyclononane-N,N',N"-triacetic acid (NOTA), 1,4,8,11-tetraazacyclotetradecane-N,N',N",N"'-tetraacetic acid (TETA) or a compound with a general formula

$$Y'''$$
 R_1 -N N- R_2
 Y'''
 Y'''
 Y'''
 Y'''

wherein

PG is a sulfur protecting group selected from alkanoyl, arylcarbonyl, arylalkanoyl, acetamidomethyl, tetrahydropyranyl and tetrahydrofuranyl,

Y', Y", and Y" are hydrogen or oxygen with the proviso that at least one of them is an O, R_1 and R_2 are hydrogen or alkyl (C_1 - C_3),

X = NH or S with the proviso that Y''' is hydrogen when X is S,

Z is PG if X is S, and

Z is hydroxyalkyl, aminoalkyl or carboxyalkyl;

 R_3 is DLys, DPhe or other D-amino acid, a spacer unit such as Gly-Gly-Gly, Gly-Ser-Gly, Tyr-Glu-Asn, DTyr-Glu-Asn, Phe-Glu-Asn or DPhe-Glu-Asn, or piperidinyl glycine (PipGly), aminomethylcyclohexylalanine (Amcha), other amino acid containing cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring;

CA is a cyclic amino acid selected from Pro, Hyp, 4-oxo-proline [4OPro], pipecolic acid (PipCA), azetidinecarboxylic acid (AzeCA), other amino acid containing cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring;

$$n = 0, 1 \text{ or } 2$$
:

 AA_1 is an amino acid containing a guanidino group except arginine, wherein the configuration at the α -carbon is either L- or D-, e.g., STLD01-918095-1

-HN
$$\alpha$$
-CO-

-HN α -

-HN α -CO-

-HN α -

-HN α

R₄ is a cycloalkyl group (C₃-C₁₀), phenyl group, aralkyl group, substituted phenyl group or substituted aralkyl group with electron withdrawing or electron donating group with the proviso that the guanidino group is present at a position not occupied by the substituent on the phenyl group;

$$p = 1-7;$$

m = 0-6;

$$q = 1-7;$$

AA₂ is arginine, lysine, piperidinylglycine (PipGly), or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-configuration at the α -carbon, or AA₂ is an amino acid containing a guanidino group wherein the configuration at the α -carbon is either L- or D-, e.g.,

AA₃ is a cyclic amino acid selected from Pro, Hyp, 4-oxo-proline [4OPro], pipecolic acid (PipCA), azetidinecarboxylic acid (AzeCA), or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D- configuration at the α -carbon;

AA₄ is Phe, Tyr, an isomer of Tyr, polyhydroxylated Phe, or other aromatic amino acid, wherein the amino acid can have the L- or D-configuration at the α -carbon;

AA₅ is t-butylglycine (tBuGly), 1-aminocyclohexylcarboxylic acid (Achc), cyclohexylglycine (Chg), trimethylsilylalanine, Ile, or other amino acid containing a branched or cyclic hydrocarbon substituent at the side chain at the α - or β -position, wherein the amino acid can have the L- or D-configuration at the α -carbon; and

AA₆ is cyclopropylalanine (Cpa), cyclohexylalanine (Cha), t-butylalanine (tBuala), Leu, or other amino acid containing a branched or cyclic hydrocarbon substituent at the side chain at the α - or β -position, wherein the amino acid can have the L- or D-configuration at the α -carbon.

Preferred compounds are:

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-Leu-OH

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-Leu-OH

DTPA-NH-CH-CO-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-Leu-OH

SEQ ID NO:4

$$DTPA-NH-CH-CO-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-Leu-OH\\$$

$$H_2NH_2C$$
 HN
 NH_2
,

SEQ ID NO:5

DTPA-DTyr-Glu-Asn-Lys-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-Leu-OH

DTPA-DTyr-Glu-Asn-Lys-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-Cha-OH

DTPA-DTyr-Glu-Asn-Lys-Pro-NH-CH-CO-Arg-Pro-Tyr-tBuGly-tBuAla-OH

DTPA-DLys-Pro-Gly(PipAm)-Arg-(4-oxo)Pro-Tyr-tBuGly-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-(2,6diMe)Tyr-tBuGly-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-mTyr-tBuGly-Leu-OH, wherein mTyr stands for meta-tyrosine such that the -OH group of the Tyr is in the meta position,

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DTPA-DLys-Pro-Gly(PipAm)-PipGly-Pro-Tyr-tBuGly-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-AzeCA-Tyr-tBuGly-Leu-OH,
DTPA-DLys-AzeCA-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Achc-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Chg-Leu-OH,
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-DTyr-tBuGly-Leu-OH,
with compounds XI and XII being the most preferred compounds.

Serum stability of these compounds was considerably increased upon substitution of AA₅ with t-butylglycine as shown in Table 3.

Table 3

		% of pe	ntide of
Sequence	K _d	% of peptide at 4 hours after incubation at 37EC	
		Serum	Urine
DTPA-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:3)	40	1.6	13.0
DTPA-DLys-Pro-Arg-(4-Gu)Phe-Pro-Tyr-Ile-Leu-OH	83	16.3	0
DTPA-DLys-Pro-(4-Gu)Phe-Arg-Pro-Tyr-Ile-Leu-OH (Compound I)	8.6	14.1	1.5
DTPA-DLys-Pro-(4-Gu)Phe-(4-Gu)Phe-Pro-Tyr-Ile-Leu-OH	175	25.2	2.3
DTPA-DLys-Pro-Arg-Aba(Apy)-Pro-Tyr-Ile-Leu-OH	1200	15.7	4.7
DTPA-DLys-Pro-Aba(Apy)-Arg-Pro-Tyr-Ile-Leu-OH	68	4.3	6.0
DTPA-DLys-Pro-Aba(Apy)-Aba(Apy)-Pro-Tyr-Ile-Leu-OH	>10000	ND	ND
DTPA-DLys-Pro-(4-Gu)Phe-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound IX)	12.5	72.0	51.1
DTPA-DLys-Pro-(4-Gu)Phe-Arg-Pro-Tyr-Leu-(Ψ-CH ₂ -NH)- Leu-OH	>10000	76.2	95.7
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Ile-Leu-OH (Compound II)	2.7	19.5	16.6
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound X)	3.5	87.5	63.0
DTPA-DLys-Pro-Gly(PipAm)-Arg-(4-oxo)Pro-Tyr-tBuGly-	>1000	78.5	79.7

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-(2,6-diMe)Tyr-tBuGly-Leu-OH 45 96.6 99.2 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-mTyr-tBuGly-Leu-OH 300 84.4 57.8 DTPA-DLys-Pro-Gly(PipAm)-Arg-ProR-OCO-Tyr-tBuGly-Leu-OH (wherein ProR-OCO indicates that the bond between the proline and the OCO is a reduced peptide bond) >10000 ND ND DTPA-DLys-Pro-Gly(PipAm)-PipGly-Pro-Tyr-tBuGly-Leu-OH 740 79.2 36.5 DTPA-DLys-Pro-Gly(PipAm)-Arg-AzeCA-Tyr-tBuGly-Leu-OH 24 79.5 49.8 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH >10000 89.9 84.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH 290 88.1 73.6 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH 12 81.2 65.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH 7.3 89.2 72.3 DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH >1000 69.5 45.8 DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH >1000 69.5 45.8	Leu-OH			
OH DTPA-DLys-Pro-Gly(PipAm)-Arg-ProR-OCO-Tyr-tBuGly-Leu-OH (wherein ProR-OCO indicates that the bond between the proline and the OCO is a reduced peptide bond) DTPA-DLys-Pro-Gly(PipAm)-PipGly-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-AzeCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-AzeCA-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH		45	96.6	99.2
Leu-OH (wherein ProR-OCO indicates that the bond between the proline and the OCO is a reduced peptide bond) DTPA-DLys-Pro-Gly(PipAm)-PipGly-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-AzeCA-Tyr-tBuGly-Leu-OH DTPA-DLys-AzeCA-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH		300	84.4	57.8
OH DTPA-DLys-Pro-Gly(PipAm)-Arg-AzeCA-Tyr-tBuGly-Leu-OH 740 79.2 36.5 OH 79.2 36.5 DTPA-DLys-AzeCA-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH 24 79.5 49.8 OH >10000 89.9 84.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH 290 88.1 73.6 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH 12 81.2 65.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH 7.3 89.2 72.3 OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH >1000 69.5 45.8 OH DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu-OH >1000 96.2 91.9	Leu-OH (wherein Pro ^R -OCO indicates that the bond between	>10000	ND	ND
OH DTPA-DLys-AzeCA-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH 24 79.5 49.8 OH >10000 89.9 84.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH 290 88.1 73.6 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH 12 81.2 65.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH 7.3 89.2 72.3 OH >1000 69.5 45.8 OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH >1000 69.5 45.8 DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu-OH >1000 69.5 45.8		140	91.6	79.7
OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Achc-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm		740	79.2	36.5
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH 290 88.1 73.6 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH 12 81.2 65.9 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH 7.3 89.2 72.3 OH 73.6 DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH 7.3 89.2 72.3 DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH 7.3 89.2 72.3 DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-7.00 89.5 45.8 DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu-7.00 96.2 91.9		24	79.5	49.8
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu->1000 69.5 45.8 DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu->1000 96.2 91.9	DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Achc-Leu-OH	>10000	89.9	84.9
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu->1000 96.2 91.9	DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH	290	88.1	73.6
OH DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu- OH DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu- >1000 96.2 91.9	DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH	12	81.2	65.9
OH DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu- >1000 96.2 91.9		7.3	89.2	72.3
		>1000	69.5	45.8
		>1000	96.2	91.9

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Chg-Leu-OH	25	75.2	7.0
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Ile ^R -OCO-Leu-OH (wherein Ile ^R -OCO indicates a reduced peptide bond between Ile and OCO)	>1000	94.6	89.8
DTPA-(Pip)Ala-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (SEQ ID NO:6) (wherein (Pip)Ala represents piperidinyl alanine)	>1000	98.9	98.3
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-DTyr-tBuGly-Leu-OH	6.4	98.9	98.3
DTPA-DLys-Pro-Ala(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (where Ala(PipAm) represents (N-amidinopiperidinyl)alanine)	14	96.5	89.8
DTPA-DLys-Pro-homoAla(PipAm)-Arg-Pro-Tyr-tBuGly- Leu-OH (wherein homoAla(PipAm) represents (N- amidinopiperidinyl)homoalanine)	14	84.3	71.3
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-HA (wherein HA indicates that this peptide ends in hydroxamic acid)	49	89.2	60.2
DTPA-PipGly-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound XI) (SEQ ID NO:4)	3.9	95.9	93.6
DTPA-trans-Cha(4-CH ₂ NH ₂)-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound XII) (SEQ ID NO:5)	5.0	94.8	89.4
DTPA-DTyr-Glu-Asn-Lys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound XIII)	4.5	94.9	86.8
DTPA-DTyr-Glu-Asn-Lys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH (Compound XIV)	4.5	93.9	81.8

DTPA-DTyr-Glu-Asn-Lys-Pro-Gly(PipAm)-Arg-Pro-Tyr-	3.5	98.9	89.8
tBuGly-tBuAla-OH (Compound XV)			

While serum stability was increased by the incorporation of Arg mimics within neurotensin, the HPLC data suggested that the C-terminal portion of the peptide may also undergo degradation (due to the presence of metabolites with slightly shorter retention times). To address this problem, additional derivatives were prepared which retained the Arg mimic that contributed to the lowest IC₅₀. The C-terminus was then modified to impart greater enzymatic stability.

	IC_{50} (nM)
DTPA-DLys-Pro-Phe(4-guanyl)-Arg-Pro-Tyr-tBuGly-Leu-OH	12.5
DTPA-DLys-Pro-Phe(4-guanyl)-Arg-Pro-Tyr-Leu(Ψ-CH ₂ NH)-Leu-NH ₂	>1000
DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH	3.5

<u>Table 4</u>

<u>Serum and Urine Stability of Neurotensin Derivatives</u>

<u>Modified at the C-Terminus</u>

Compound	Prep	Serum (t=4h)	Urine (t=4h)
In-111-DTPA-R-R-P-Y-I-L (SEQ ID NO:3)	97.6%	1.6%	14.0%
	rt=14.2m	rt=14.2m	rt=14.2m
In-111-DTPA-DK-P-F(Gu)-R-P-Y-I-L	98.7%	14.1%	1.5%
	rt=14.6m	rt=14.6m	rt=14.6m
In-111-DTPA-DK-P-G(PipAm)-R-P-Y-I-L	99.3%	19.5%	16.6%
	rt=13.8m	rt=13.8m	rt=13.8m

In-111-DTPA-DK-P-F(Gu)-R-P-Y-tBuG-L	99.0%	72.0%	51.1%
	rt=13.6	rt=13.6m	rt=13.6m
In-111-DTPA-DK-P-F(Gu)-R-P-Y-L(Ψ-CH ₂ NH)-L-	99.7%	76.2%	95.7%
NH ₂	rt=13.3m	rt=13.3m	rt=13.3m
In-111-DTPA-DK-P-G(PipAm)-R-P-Y-tBuG-L	99.2%	87.5%	63.0%
	rt=13.8m	rt=13.8m	rt=13.8m

While the invention has been disclosed in this patent application by reference to the details of preferred embodiments of the invention, it is to be understood that the disclosure is intended in an illustrative rather than in a limiting sense, as it is contemplated that modifications will readily occur to those skilled in the art, within the spirit of the invention and the scope of the appended claims.

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Mazella J, et al. (1983) J. Biol. Chem. 258:3476-3481.

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[WHAT IS CLAIMED IS:] CLAIMS

1. A peptide of structure CM-R₃-(CA)_n-AA₁-AA₂-AA₃-AA₄-AA₅-AA₆-OH, wherein said peptide has a selective affinity for neurotensin receptors and wherein

CM is a chelating moiety or metal binding site;

 R_3 is D-lysine, D-phenylalanine, any D-amino acid, glycine-glycine-glycine, Gly-Ser-Gly, Tyr-Glu-Asn, DTyr-Glu-Asn, Phe-Glu-Asn, DPhe-Glu-Asn, piperidinyl glycine, aminomethylcyclohexylalanine, amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring, or a spacer unit;

CA is a cyclic amino acid selected from the group consisting of proline, hydroxyproline, 4-oxo-proline, pipecolic acid, azetidinecarboxylic acid, and other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring;

n = 0, 1 or 2;

 AA_1 is an amino acid which comprises a guanidino group and wherein the α -carbon is either L- or D-, with the proviso that AA_1 is not arginine;

AA₂ is arginine, lysine, piperidinylglycine, or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-

configuration at the α -carbon, or AA_2 is an amino acid which comprises a guanidino group wherein the α -carbon is either L- or D-;

AA₃ is a cyclic amino acid selected from proline, hydroxyproline, 4-oxo-proline, pipecolic acid, azetidinecarboxylic acid, or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-configuration at the α -carbon;

AA₄ is phenylalanine, tyrosine, an isomer of tyrosine, polyhydroxylated phenylalanine, or other aromatic amino acid, wherein the amino acid can have the L- or D-configuration at the α -carbon;

AA₅ is isoleucine; and

AA₆ is leucine.

2. The peptide of claim 1 wherein AA_1 is

wherein

m = 0-6;

p = 1-7;

q = 1-7; and

R₄ is cycloalkyl C₃-C₁₀, phenyl, aralkyl, substituted phenyl or substituted aralkyl comprising an electron withdrawing or electron donating group with the proviso that said

guanidino group is at a position different from said electron withdrawing or electron donating group.

- 3. The peptide of claim 1 wherein said peptide is labeled with a radioisotope.
- 4. The peptide of claim 3 wherein said label is ^{99m}Tc, ²⁰³Pb, ⁶⁷Ga, ¹¹¹In, ⁹⁷Ru, ⁶²Cu, ⁶⁴Cu, ¹⁸⁶Re, ¹⁸⁸Re, ⁹⁰Y, ¹²¹Sn, ¹⁶¹Tb, ¹⁵³Sm, ¹⁶⁶Ho, ¹⁰⁵Rh, ¹⁷⁷Lu or a radioactive halogen isotope.
- 5. The peptide of claim 4 wherein if said label is a metal then CM is a chelating group for said metal and if said label is a halogen then said halogen is bound to an aromatic ring.
- 6. The peptide of claim 1 wherein CM is ethylene diamine tetraacetic acid (EDTA), diethylene triamine pentaacetic acid (DTPA), cyclohexyl 1,2-diamine tetraacetic acid (CDTA), ethyleneglycol-O,O=-bis(2-aminoethyl)-N,N,N',N'-diacetic acid (HBED), triethylene tetraamine hexaacetic acid (TTHA), 1,4,7,10-tetraazacyclododecane-N,N',N",N"'-tetraacetic acid (DOTA), 1,4,7-triazacyclononane-N,N',N"-triacetic acid (NOTA), 1,4,8,11-tetraazacyclotetradecane-N,N',N",N"'-tetraacetic acid (TETA) or a compound of formula

wherein

PG is a sulfur protecting group selected from the group consisting of alkanoyl, arylcarbonyl, arylalkanoyl, acetamidomethyl, tetrahydropyranyl and tetrahydrofuranyl; Y', Y'', and Y''' are hydrogen or oxygen with the proviso that at least one of them is an O; R_1 and R_2 are hydrogen or alkyl (C_1 - C_3);

X = NH or S with the proviso that Y''' is hydrogen when X is S;

Z is PG if X is S; and

Z is hydroxyalkyl, aminoalkyl or carboxyalkyl.

7. The peptide of claim 1 wherein said peptide is

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

$$N$$
 H_2N
III

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

$$V$$
 N
 NH_2

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

DTPA-DLys-Pro-NH-CH-CO-Arg-Pro-Tyr-Ile-Leu-OH

or

A peptide of structure CM-R₃-(CA)_n-AA₁-AA₂-AA₃-AA₄-AA₅-AA₆-OH, wherein said peptide has a selective affinity for neurotensin receptors and wherein CM is a chelating moiety or metal binding site;

 R_3 is D-lysine, D-phenylalanine, any D-amino acid, glycine-glycine-glycine, Gly-Ser-Gly, Tyr-Glu-Asn, DTyr-Glu-Asn, Phe-Glu-Asn, DPhe-Glu-Asn, piperidinyl glycine, aminomethylcyclohexylalanine, other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring, or a spacer unit;

CA is a cyclic amino acid selected from the group consisting of proline, hydroxyproline, 4-oxo-proline, pipecolic acid, azetidinecarboxylic acid, other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or an alkyl amino substituent either externally or as a part of the ring;

n = 0, 1 or 2;

 AA_1 is an amino acid which comprises a guanidino group and wherein the α -carbon is either L- or D-, with the proviso that AA_1 is not arginine;

AA₂ is arginine, lysine, piperidinylglycine, or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-configuration at the α -carbon, or AA₂ is an amino acid which comprises a guanidino group wherein the α -carbon is either L- or D-;

AA₃ is proline, hydroxyproline, 4-oxo-proline, pipecolic acid, azetidinecarboxylic acid, or other amino acid containing a cycloalkyl ring at the α - or β -position with an amine group or alkyl amino substituent either externally or as a part of the ring, wherein the amino acid can have the L- or D-configuration at the α -carbon;

AA₄ is phenylalanine, tyrosine, an isomer of tyrosine, polyhydroxylated phenylalanine, or other aromatic amino acid wherein said amino acid can have the L- or D-configuration at the α -carbon;

AA₅ is t-butylglycine, 1-aminocyclohexylcarboxylic acid, cyclohexylglycine, trimethylsilylalanine, isoleucine, or other amino acid containing a branched or cyclic hydrocarbon substituent at the side chain at the α - or β -position, wherein the amino acid can have the L- or D-configuration at the α -carbon; and

AA₆ is cyclopropylalanine, cyclohexylalanine, t-butylalanine, leucine, or other amino acid containing a branched or cyclic hydrocarbon substituent at the side chain at the α - or β -position, wherein the amino acid can have the L- or D-configuration at the α -carbon.

9. The peptide of claim 8 wherein AA₁ is

$$m = 0-6;$$

 $p = 1-7;$

q = 1-7; and

R₄ is cycloalkyl C₃-C₁₀, phenyl, aralkyl, substituted phenyl or substituted aralkyl comprising an electron withdrawing or electron donating group with the proviso that said guanidino group is at a position different from said electron withdrawing or electron donating group.

- 10. The peptide of claim 8 wherein said peptide is labeled with a radioisotope.
- 11. The peptide of claim 10 wherein said label is ^{99m}Tc, ²⁰³Pb, ⁶⁷Ga, ¹¹¹In, ⁹⁷Ru, ⁶²Cu, ⁶⁴Cu, ¹⁸⁶Re, ¹⁸⁸Re, ⁹⁰Y, ¹²¹Sn, ¹⁶¹Tb, ¹⁵³Sm, ¹⁶⁶Ho, ¹⁰⁵Rh, ¹⁷⁷Lu or a radioactive halogen isotope.
- 12. The peptide of claim 11 wherein if said label is a metal then CM is a chelating group for said metal and if said label is a halogen then said halogen is bound to an aromatic ring.

The peptide of claim 8 wherein CM is ethylene diamine tetraacetic acid (EDTA), diethylene triamine pentaacetic acid (DTPA), cyclohexyl 1,2-diamine tetraacetic acid (CDTA), ethyleneglycol-O,O=-bis(2-aminoethyl)-N,N,N',N'-diacetic acid (HBED), triethylene tetraamine hexaacetic acid (TTHA), 1,4,7,10-tetraazacyclododecane-N,N',N'',N'''-tetraacetic acid (DOTA), 1,4,7-triazacyclononane-N,N',N''-triacetic acid (NOTA), 1,4,8,11-tetraazacyclotetradecane-N,N',N'',N'''-tetraacetic acid (TETA) or a compound of formula

$$Y = \begin{bmatrix} Y'' \\ R_1-N & N-R_2 \\ S & X \\ PG & Z \end{bmatrix}$$

wherein

PG is a sulfur protecting group selected from the group consisting of alkanoyl, arylcarbonyl, arylalkanoyl, acetamidomethyl, tetrahydropyranyl and tetrahydrofuranyl; Y', Y'', and Y''' are hydrogen or oxygen with the proviso that at least one of them is an O; R_1 and R_2 are hydrogen or alkyl (C_1 - C_3);

X = NH or S with the proviso that Y''' is hydrogen when X is S;

Z is PG if X is S, and

Z is hydroxyalkyl, aminoalkyl or carboxyalkyl.

14. The peptide of claim 8 wherein said peptide is DTPA-Arg-Arg-Pro-Tyr-Ile-Leu-OH (SEQ ID NO:3), DTPA-DLys-Pro-Arg-(4-Gu)Phe-Pro-Tyr-Ile-Leu-OH, DTPA-DLys-Pro-(4-Gu)Phe-Arg-Pro-Tyr-Ile-Leu-OH (Compound I), DTPA-DLys-Pro-(4-Gu)Phe-(4-Gu)Phe-Pro-Tyr-Ile-Leu-OH,

DTPA-DLys-Pro-Arg-Aba(Apy)-Pro-Tyr-Ile-Leu-OH,

DTPA-DLys-Pro-Aba(Apy)-Arg-Pro-Tyr-Ile-Leu-OH,

DTPA-DLys-Pro-Aba(Apy)-Aba(Apy)-Pro-Tyr-Ile-Leu-OH,

DTPA-DLys-Pro-(4-Gu)Phe-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound IX),

DTPA-DLys-Pro-(4-Gu)Phe-Arg-Pro-Tyr-Leu(Ψ-CH₂-NH)Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Ile-Leu-OH (Compound II),

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound X),

DTPA-DLys-Pro-Gly(PipAm)-Arg-(4-oxo)Pro-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-(2,6diMe)Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-mTyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro^R-OCO-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-PipGly-Pro-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-AzeCA-Tyr-tBuGly-Leu-OH,

DTPA-DLys-AzeCA-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Achc-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cpa-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-PipCA-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-DPipCA-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Chg-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-Ile^R-OCO-Leu-OH,

DTPA-(Pip)Ala-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (SEQ ID NO:6),

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-DTyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Ala(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-homoAla(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH,

DTPA-DLys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-HA,

DTPA-PipGly-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound XI) (SEQ ID NO:4),

DTPA-trans-Cha(4-CH₂NH₂)-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound XII) (SEQ ID NO:5),

DTPA-DTyr-Glu-Asn-Lys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Leu-OH (Compound XIII),

DTPA-DTyr-Glu-Asn-Lys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-Cha-OH (Compound XIV), or

DTPA-DTyr-Glu-Asn-Lys-Pro-Gly(PipAm)-Arg-Pro-Tyr-tBuGly-tBuAla-OH (Compound XV).

- 15. A method for diagnosing a patient for a tumor by administering an effective amount of a peptide of claim 1.
- 16. The method of claim 15 wherein said tumor is a small cell lung carcinoma, exocrine pancreatic cancer, Ewing sarcoma, meningioma, medulloblastoma, or astrocytoma.
- 17. A method for diagnosing a patient for a tumor by administering an effective amount of a peptide of claim 8.
- 18. The method of claim 17 wherein said tumor is a small cell lung carcinoma, exocrine pancreatic cancer, Ewing sarcoma, meningioma, medulloblastoma, or astrocytoma.
- 19. A method for treating a patient for a tumor by administering an effective amount of a peptide of claim 1.
- 20. The method of claim 19 wherein said tumor is a small cell lung carcinoma, exocrine pancreatic cancer, Ewing sarcoma, meningioma, medulloblastoma, or astrocytoma.

- 21. A method for treating a patient for a tumor by administering an effective amount of a peptide of claim 8.
- 22. The method of claim 21 wherein said tumor is a small cell lung carcinoma, exocrine pancreatic cancer, Ewing sarcoma, meningioma, medulloblastoma, or astrocytoma.

[ABSTRACT OF THE DISCLOSURE] ABSTRACT OF THE DISCLOSURE

Peptide analogs of neurotensin are disclosed which are resistant to enzymatic degradation and which retain high binding affinity for neurotensin receptors. Pharmaceutical compositions of these compounds are useful for diagnostic and therapeutic purposes.